

024455pcus\_ST25 (2).txt  
SEQUENCE LISTING

<110> Shanghai Genomics, Inc.  
<120> TUMOR TAG AND THE USE THEREOF  
<130> 186353/US  
<140> 10/527,257  
<141> 2005-03-09  
<150> PCT/CN2002/000631  
<151> 2002-09-09  
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<212> DNA  
<213> Homo sapiens



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ctt tgc tat gac atc acc gtc atc cct aag ttc aga cct gga cca cgg 144  
Leu Cys Tyr Asp Ile Thr Val Ile Pro Lys Phe Arg Pro Gly Pro Arg  
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tgg tgt gcg gtt caa ggc cag gtg gat gaa aag act ttt ctt cac tat 192  
Trp Cys Ala Val Gln Gly Gln Val Asp Glu Lys Thr Phe Leu His Tyr  
50 55 60  
gac tgt ggc agc aag aca gtc aca ccc gtc agt ccc ctg ggg aag aaa 240  
Asp Cys Gly Ser Lys Thr Val Thr Pro Val Ser Pro Leu Gly Lys Lys  
65 70 75 80  
cta aat gtc aca acg gcc tgg aaa gca cag aac cca gta ctg aga gag 288  
Leu Asn Val Thr Thr Ala Trp Lys Ala Gln Asn Pro Val Leu Arg Glu  
85 90 95  
gtg gtg gac ata ctt aca gag caa ctg ctt gac att cag ctg gag aat 336  
Val Val Asp Ile Leu Thr Glu Gln Leu Leu Asp Ile Gln Leu Glu Asn  
100 105 110  
tac ata ccc aag gaa ccc ctc acc ctg cag gcc agg atg tct tgt gag 384  
Tyr Ile Pro Lys Glu Pro Leu Thr Leu Gln Ala Arg Met Ser Cys Glu  
115 120 125  
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Gln Lys Ala Glu Gly His Gly Ser Gly Ser Trp Gln Leu Ser Phe Asp  
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 gtt cat cct gga gcc aga aag atg aaa gaa aag tgg gag aat gac aag 528  
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 165 170 175  
 gat atg acc atg tcc ttc cat tac atc tca atg gga gac tgc aca gga 576  
 Asp Met Thr Met Ser Phe His Tyr Ile Ser Met Gly Asp Cys Thr Gly  
 180 185 190  
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 Leu Cys Tyr Asp Ile Thr Val Ile Pro Lys Phe Arg Pro Gly Pro Arg  
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 Trp Cys Ala Val Gln Gly Gln Val Asp Glu Lys Thr Phe Leu His Tyr  
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 Asp Cys Gly Ser Lys Thr Val Thr Pro Val Ser Pro Leu Gly Lys Lys  
 65 70 75 80  
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 85 90 95  
 Val Val Asp Ile Leu Thr Glu Gln Leu Leu Asp Ile Gln Leu Glu Asn  
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 Tyr Ile Pro Lys Glu Pro Leu Thr Leu Gln Ala Arg Met Ser Cys Glu  
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Gln Lys Ala Glu Gly His Gly Ser Gly Ser Trp Gln Leu Ser Phe Asp  
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Gly Gln Ile Phe Leu Leu Phe Asp Ser Glu Asn Arg Met Trp Thr Thr  
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Val His Pro Gly Ala Arg Lys Met Lys Glu Lys Trp Glu Asn Asp Lys  
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Leu Cys Tyr Asp Ile Thr Val Ile Pro Lys Phe Arg Pro Gly Pro Arg  
 35 40 45

Trp Cys Ala Val Gln Gly Gln Val Asp Glu Lys Thr Phe Leu His Tyr  
 50 55 60

Asp Cys Gly Asn Lys Thr Val Thr Pro Val Ser Pro Leu Gly Lys Lys  
 65 70 75 80

Leu Asn Val Thr Thr Ala Trp Lys Ala Gln Asn Pro Val Leu Arg Glu  
 85 90 95

Val Val Asp Ile Leu Thr Glu Gln Leu Arg Asp Ile Gln Leu Glu Asn  
 100 105 110

Tyr Thr Pro Lys Glu Pro Leu Thr Leu Gln Ala Arg Met Ser Cys Glu  
 115 120 125

Gln Lys Ala Glu Gly His Ser Ser Gly Ser Trp Gln Phe Ser Phe Asp  
 130 135 140

Gly Gln Ile Phe Leu Leu Phe Asp Ser Glu Lys Arg Met Trp Thr Thr  
 145 150 155 160

Val His Pro Gly Ala Arg Lys Met Lys Glu Lys Trp Glu Asn Asp Lys  
 165 170 175

Val Val Ala Met Ser Phe His Tyr Phe Ser Met Gly Asp Cys Ile Gly  
 180 185 190

Trp Leu Glu Asp Phe Leu Met Gly Met Asp Ser Thr Leu Glu Pro Ser  
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Ala Gly Ala Pro Leu Ala Met Ser Ser Gly Thr Thr Gln Leu Arg Ala  
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